

FIG. 1

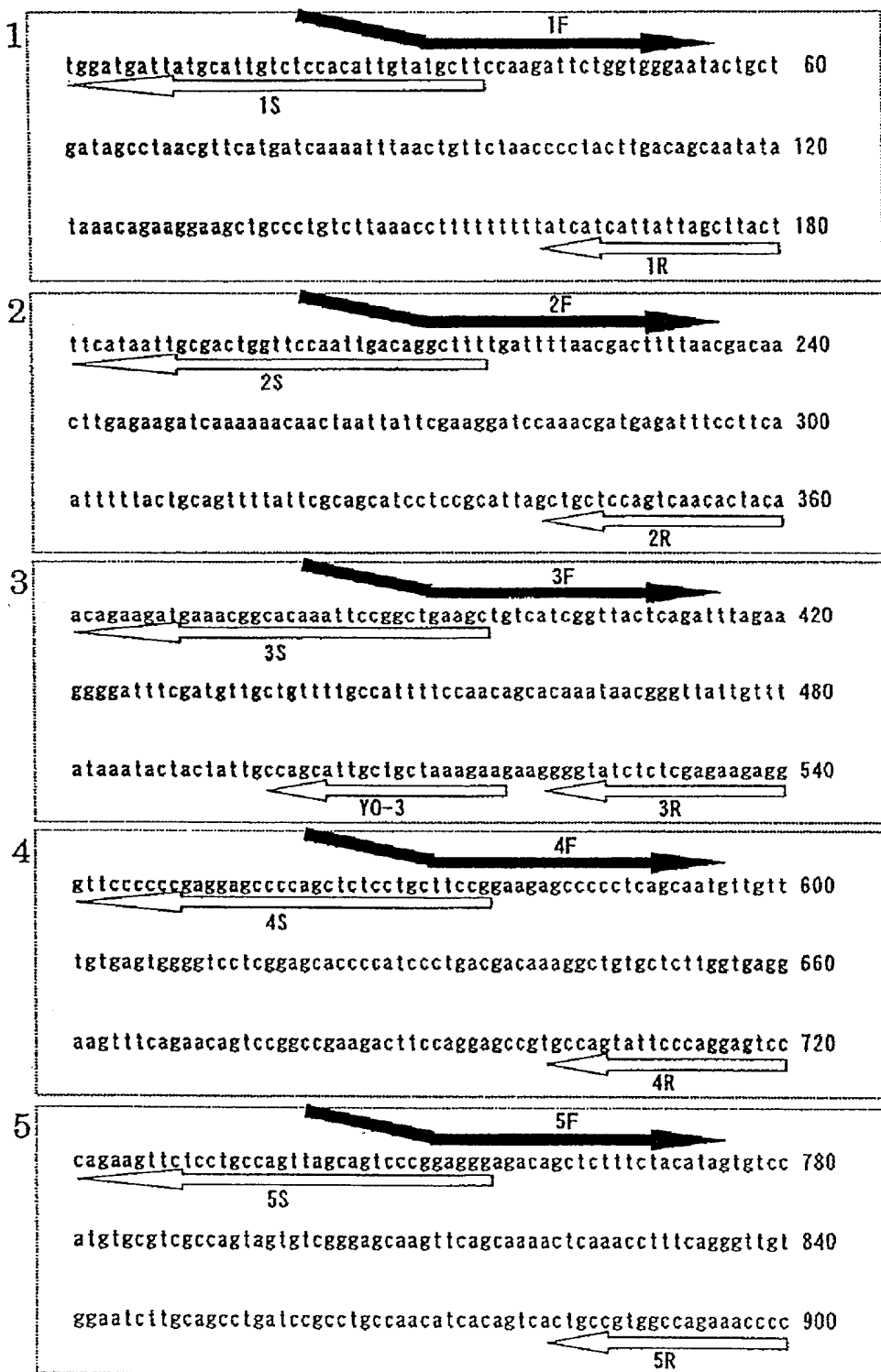


FIG. 2

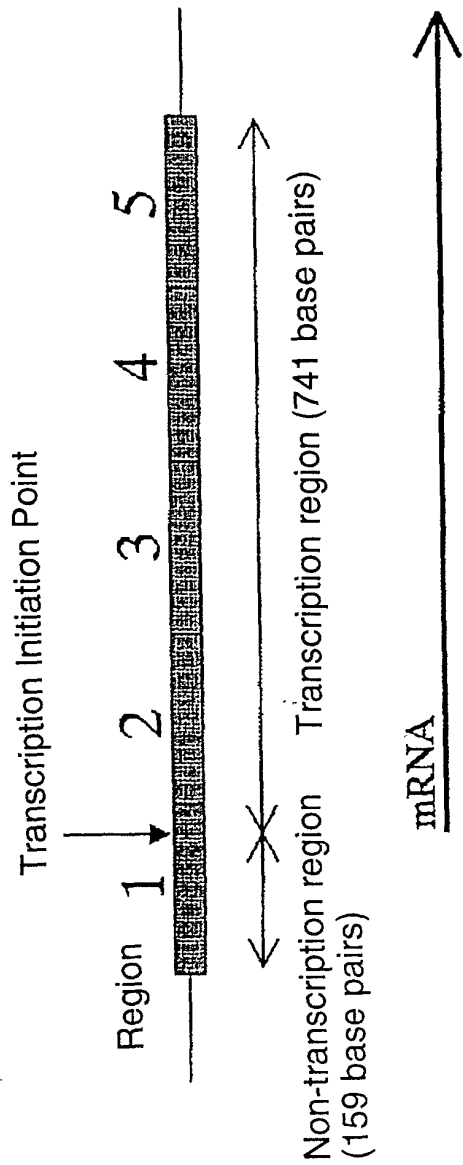
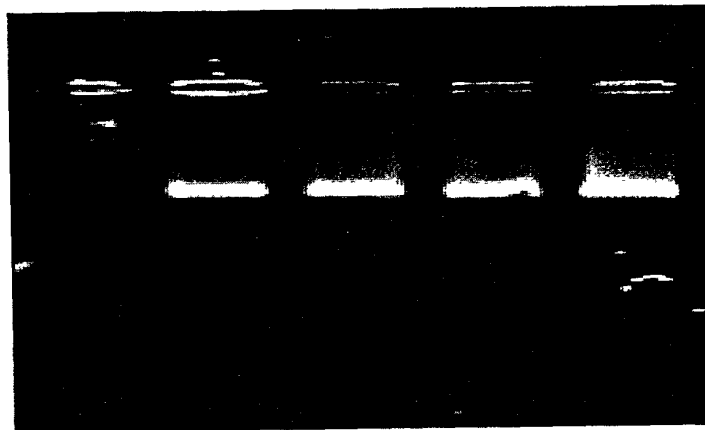


FIG. 3

(RT-PCR 30 cycles)

Specific region

1 2 3 4 5



← 178

FIG. 4A

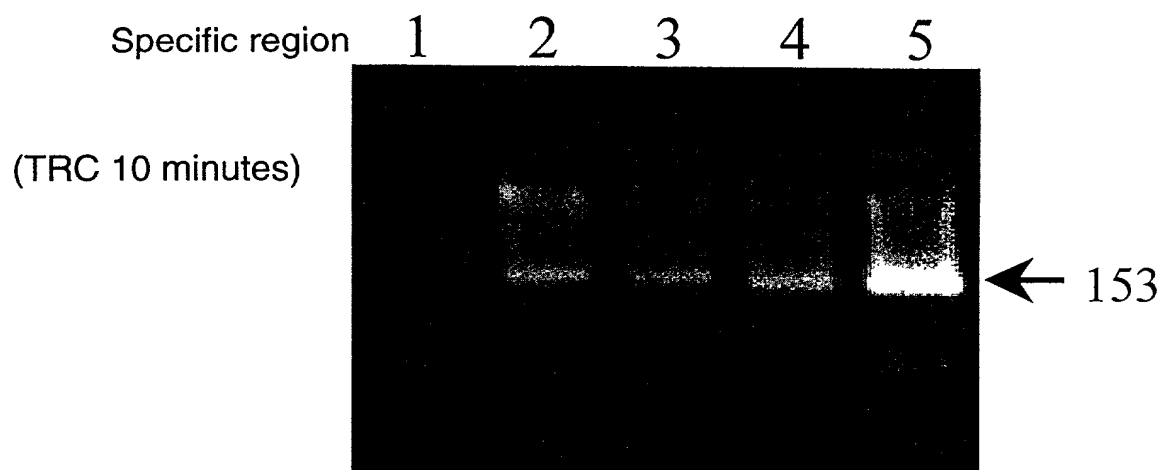


FIG. 4B

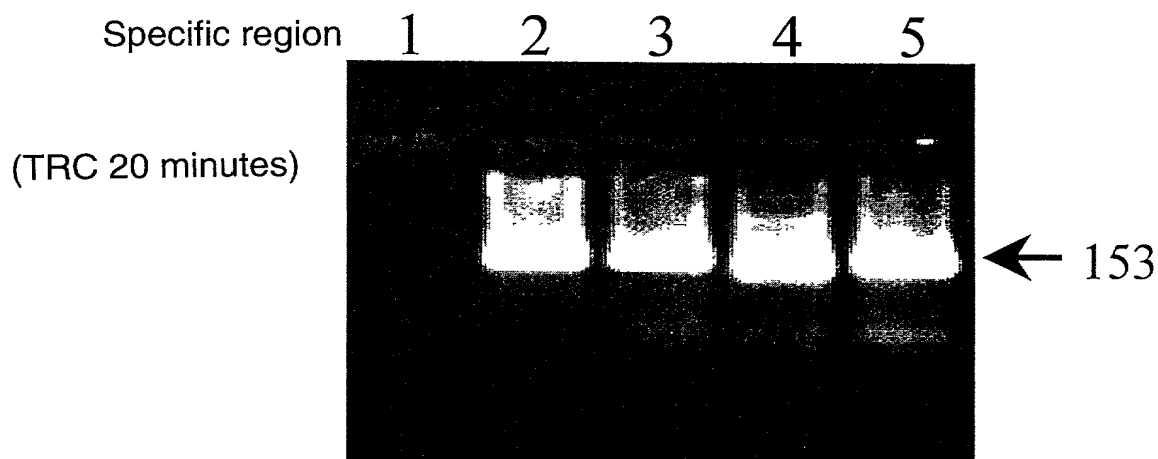


FIG. 4C

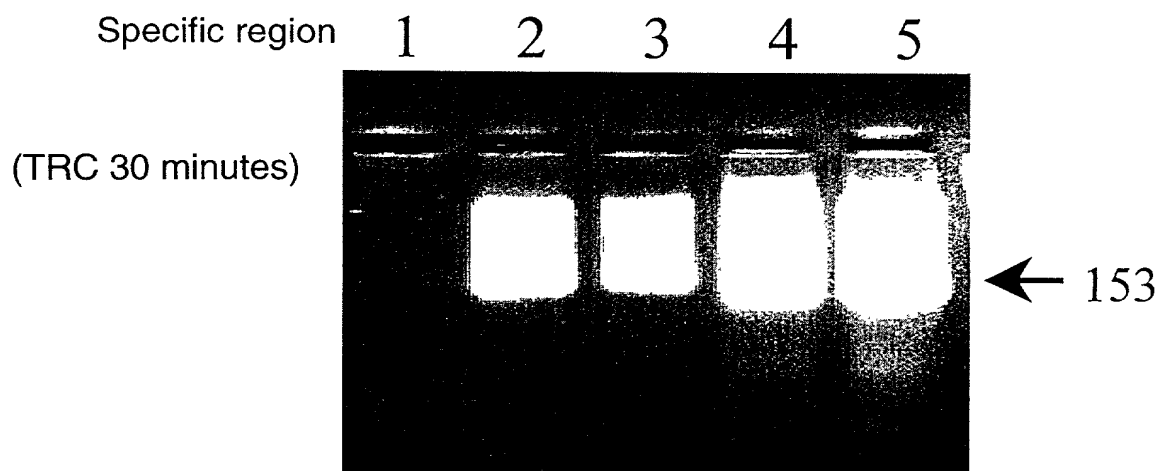


FIG. 5

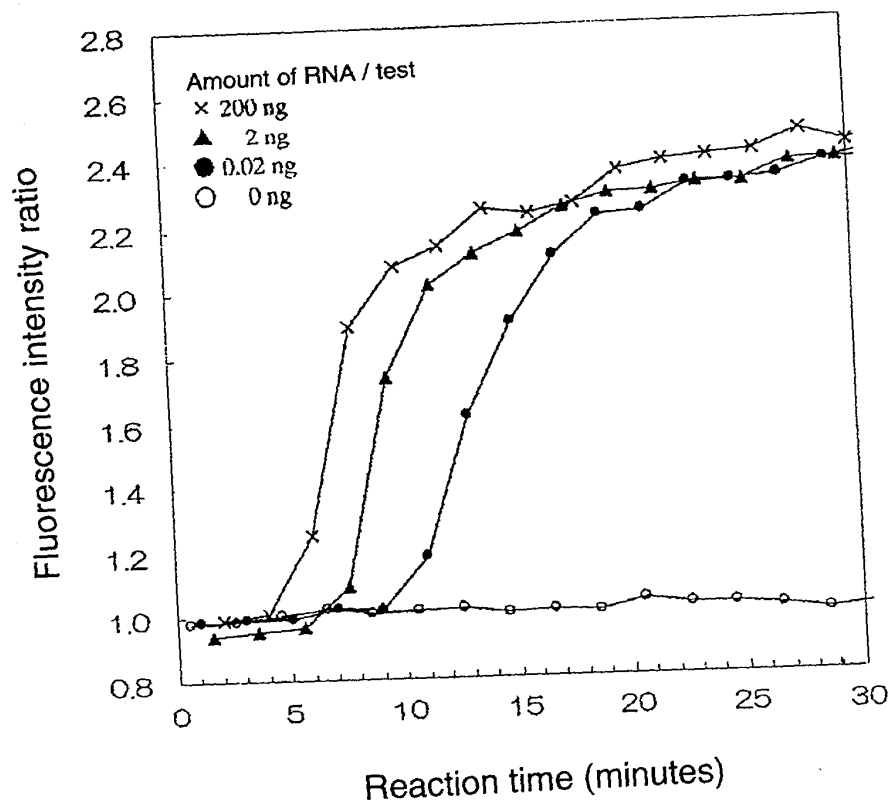


FIG. 6A

(RT-PCR 30 cycles, specific region 3)

200 20 2 0.2 0.02 0.002 0 RNA (ng)

178 →

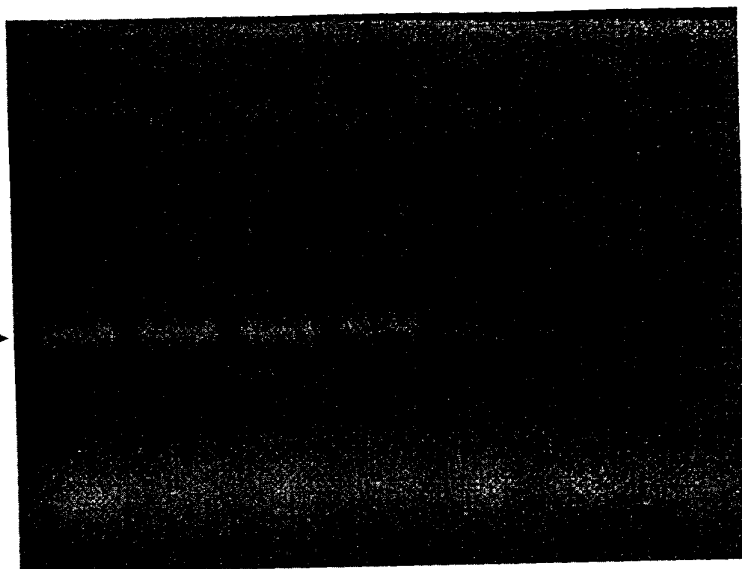


FIG. 6B

(TRC 30 minutes, specific region 3)

200 20 2 0.2 0.02 0.002 0 RNA (ng)

153 →

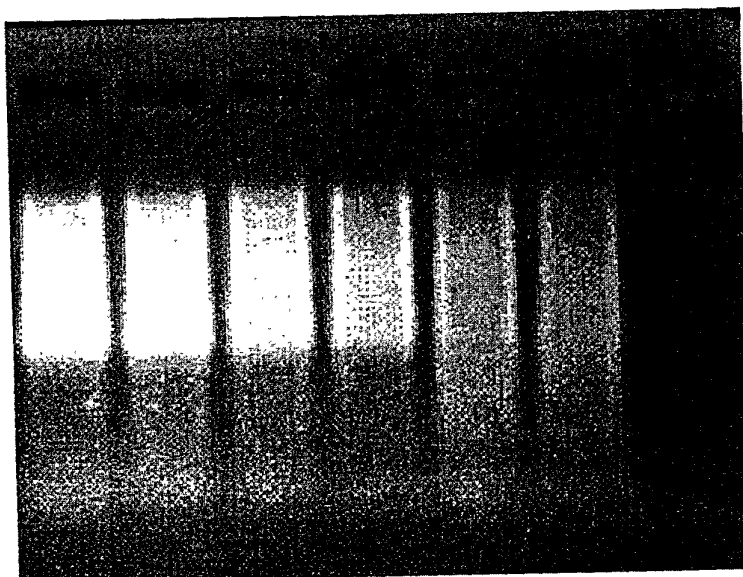


FIG. 7A

(RT-PCR 30 cycles, specific region 3)

2	0	0	0	0	mRNA (ng)
0	0	2	20	200	DNA (ng)

178 →

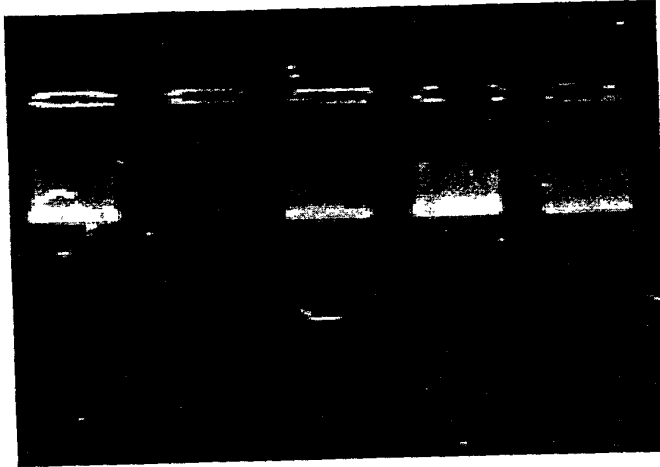


FIG. 7B

(TRC 30 minutes, specific region 3)

2	0	0	0	0	mRNA (ng)
0	0	2	20	200	DNA (ng)

153 →

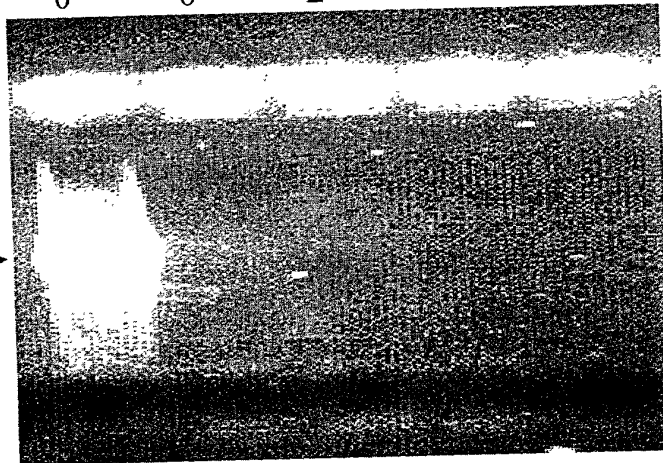


FIG. 8

